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Figure 1

Complete nucleotide sequence of the cDNA of the protein Ki-67 and the protein amino acid sequence derived therefrom.

Sequence data for the cDNA of the protein Ki-67 and the derived protein amino acid sequence. The sequence is presented in two columns, with the first column containing the nucleotide sequence and the second column containing the corresponding amino acid sequence. The sequence starts with a start codon (ATG) at position 1 and ends with a stop codon (TAA) at position 223. The sequence includes several exons, with the first exon spanning from position 1 to 135, the second exon spanning from position 136 to 200, and the third exon spanning from position 201 to 223. The sequence also includes a poly-A tail at the 3' end. The derived protein amino acid sequence is shown in bold. The sequence includes a nuclear targeting sequence (KDEL) at positions 186-190 and a nuclear targeting sequence (KDEL) at positions 210-214.

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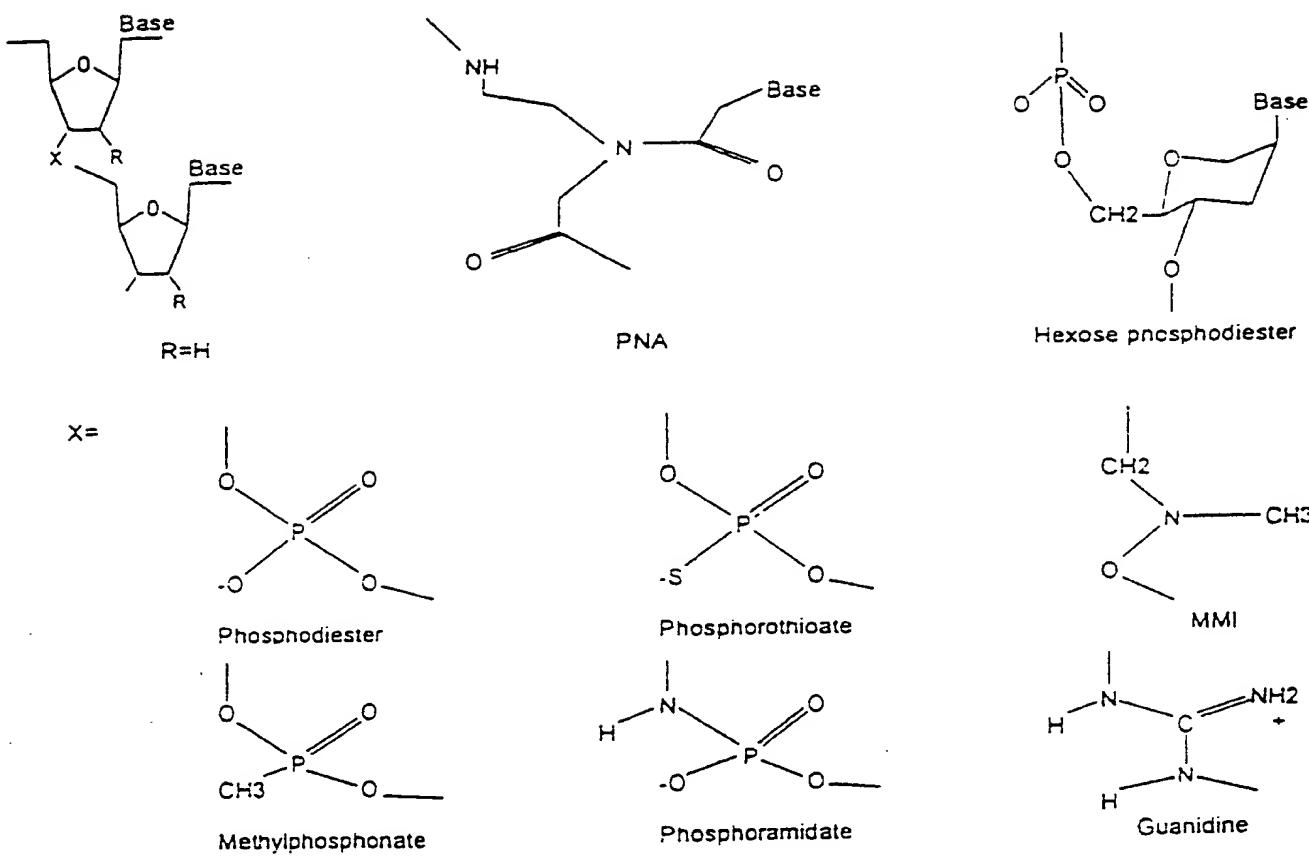
Figure 1
(continued)

TCTAAAGATATTCACAGAGTCTCCGGGCGCTTAACTAGAACCCCTGGGAGACUGGCT 9120
 L K E S A K V L R A D K V S P V S D V V 2975
 AUUCGCCAGAGACCCCTGAAATACACAAACCAAGGAACTTCTCTGGCCCACTGGCC 9180
 S T R D > V K E D S K S R T C S I F P I Y 2975
 CTCAGACGGGAACTGGCAACATGGAAACGGTACGGGAAACAGAGGGCTGGCCCAT 9240
 P X R G G C Y D G E V T G T K R C R 3025
 [---BIPARTITE NUCLEAR TARGETING SEQUENCE---]
 UCCAUACACACAGAAATTGGGAGGACCTGGCAACGAAACAGAGGGCTGGCC 9300
 P A P E S I V E E S I F P A S K R G K V A 3025
 TAGGCTAGAGGCAAATGATCCAAACCCCTGGCTCATCATGAAAGACAACTTGGGCTT 9360
 P A R S X S S E P V V C H R F S L H T S 3055
 [---] AT/CTG-BINDING MOTIF & Y-LOOP
 TCCAAACACAACTGGCAACGAACTGGCAACGAAACACATCAAACCAACAGAA 9420
 A F R E S E P A Z E L N S H D N K T H R E 3075
 CGAACACAAAATACAGACTGGCTCTGGAAATAACCCAAATACCTGGCTGGAGG 9480
 E B Y D O S V P S S F C I S L E D R R 3095
 END OF THE LARGE EXON 13 [---]
 CGAAGATACACTGGCAACGAAATAACTGGCAACGCTCTCTTATTAGCAAGAGAA 9540
 Q D K T V C A Z C O S I T V C V V L A S R I 3115
 AGAAATACACAGAAATGAAAGAAACCCATGAGACCTTCCACAGATGAGAACTCACAA 9600
 E I P R E K D P R Y T R F C E D I C R 3135
 TCCACAGATGGACCGGAAACCCATACTTACAGACAAAGCTACTGAAACAAAGG 9660
 P D D G A F K F I Z P K D V T I K F F C 3135
 [---BIPARTITE NUCLEAR TARGETING SEQUENCE---]
 CTGAGGCTCTCTACAGAGAAATGAGGCTCCACGCTTACGCTGAGGAGAGGG 9720
 I R S A X Q H V S S S P D V A E E S S G C 3175
 CGACACACTCCGAGGTTTATGCAAAATAGAAGGGGAGACGAGGAA 9780
 S Y S A X V L K H O D K G Y G E S A G H S 3195
 AGACTCCATCTGGGATGAAAGAAAGACAAACGGCACTTCCACGAGGAACTTGG 9840
 P E K C I R S R R T K S C P A A E T L Z 3215
 GGGAAATCTCCACACAGTACGGGCACTGTCAGGGTGTGCAAAATCCAGGAA 9900
 S R S V Q R V T R S V F R C S E H T F X 3235
 GCGTACCCACAACTGCTGAAAGAAATAACACAACTGAGCTTAAAGGACAGTCAGAA 9950
 A F D N V C V X X E T T S R E D S E D 3255
 TATTGACAGAAAAATCCAACTGGGAAATAATAATAATTAGTTTTCAGAGTCT 10020
 3255
 ATGCGAGTTTCTATTAACAGGAACTTCTGAGTAACTTCTAGGCTGCTTA 10080
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 CTGCTGGCTTCTGGACGAGAACTTCTCACTGGCTCTGAAAGCTTAAATCTCACTGG 10200
 CAGGTTTCTGGCTTCTGGACGAGAACTTCTCACTGGCTCTGAAAGCTTAAATCTCACTGG 10260
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 AGGCTTCTGGCTTCTGGAGAGACTTCTCACTGGCTCTGAAAGCTTAAATCTCACTGG 11580
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 GCGCTGGCTTCTGGAGAGACTTCTCACTGGCTCTGAAAGCTTAAATCTCACTGG 11820
 CCCCCGTTCTGGCTTCTGGAGAGACTTCTCACTGGCTCTGAAAGCTTAAATCTCACTGG 11880
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 ACCTTCTGGCTTCTGGAGAGACTTCTCACTGGCTCTGAAAGCTTAAATCTCACTGG 12240
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 GTATTTCTGGCTTCTGGAGAGACTTCTCACTGGCTCTGAAAGCTTAAATCTCACTGG 12480
 ATGAAATCTGGCTTCTGGAGAGACTTCTCACTGGCTCTGAAAGCTTAAATCTCACTGG 12540

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Figure 2

Structure of sugar- and phosphate-modified oligonucleotides



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Figure 3

Influence of oligonucleotides on RT4 cells.

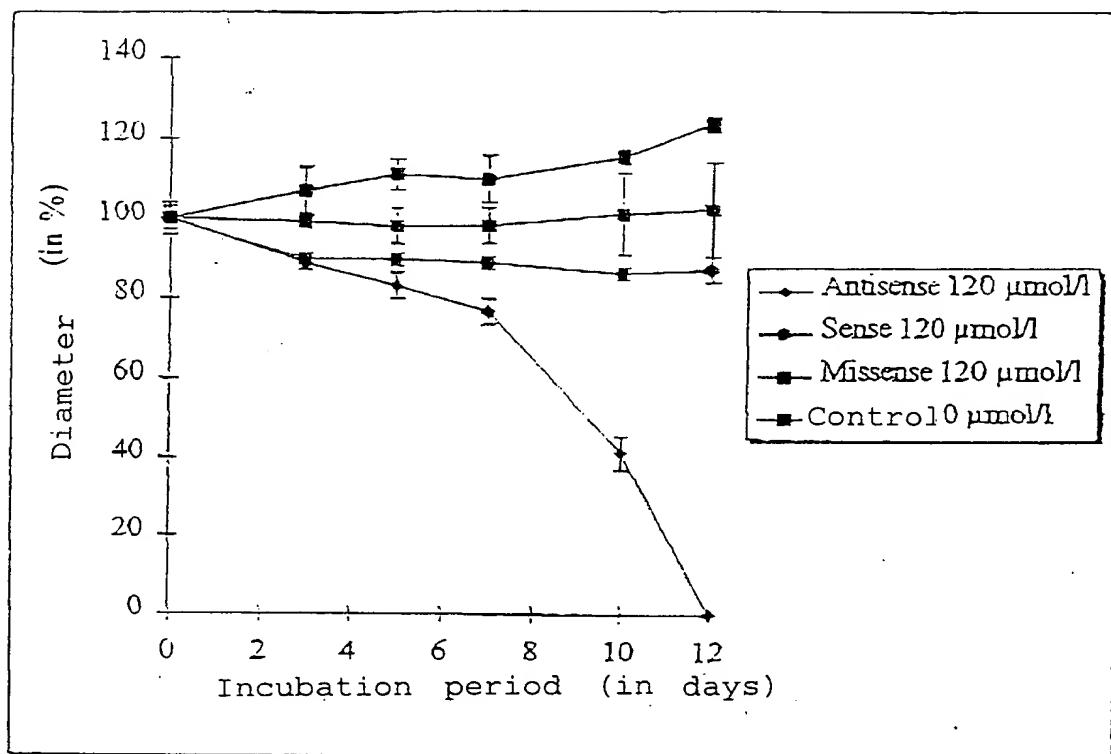
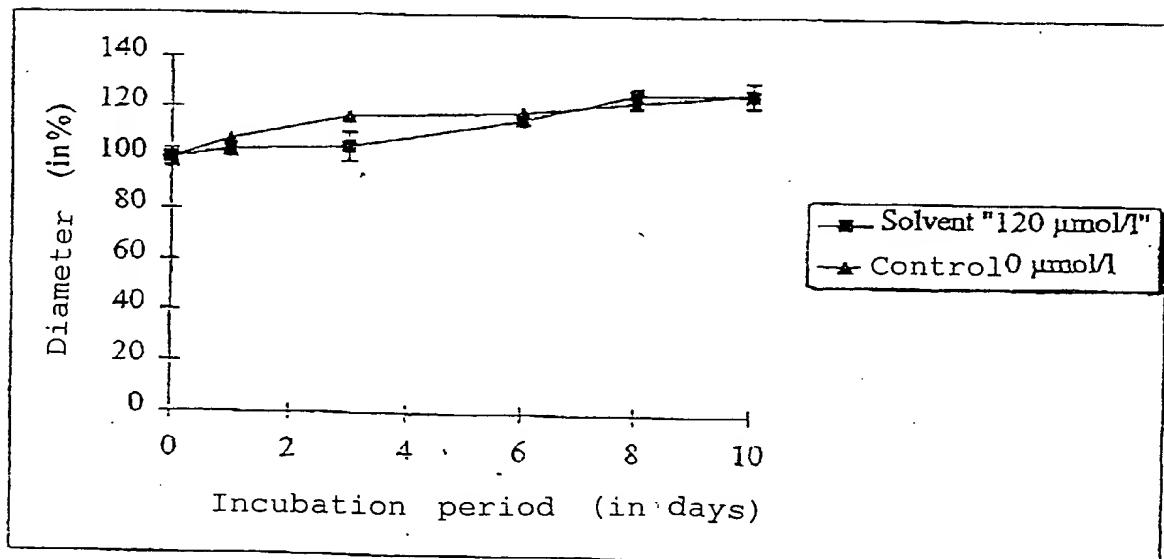


Figure 4

Influence of the solvent on RT4 cells
(negative control)

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09700906 . 022601
09/700906

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Figure 5

Influence of oligonucleotides on RT4 cells by microinjection

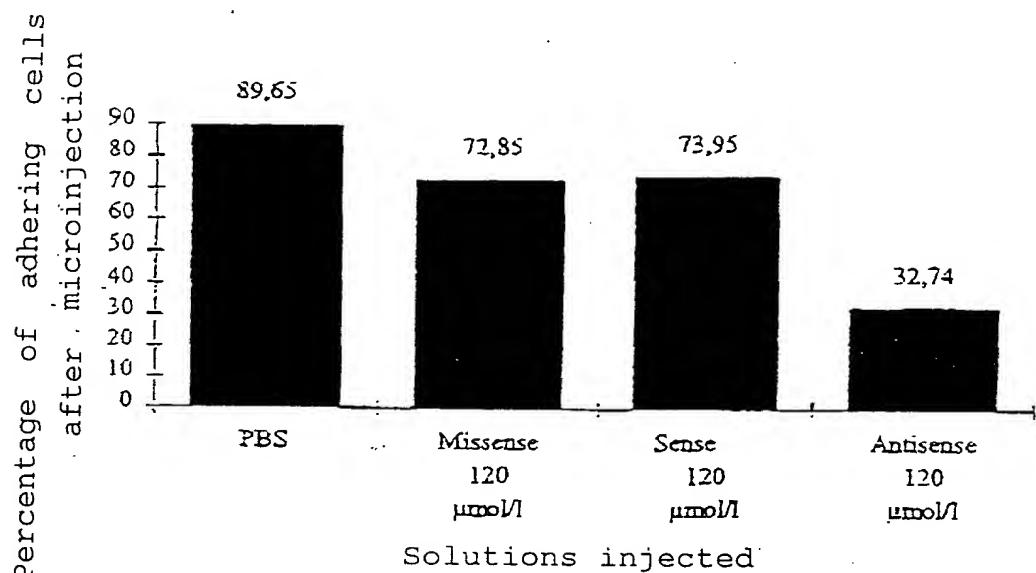


Figure 6

Influence of oligonucleotides on J82 cells

